

Nickol

Serial Number:

09/580,803

CRF Processing Date:

2/17/2001

Edited by:

Verified by:

(STIC staff)

ENTERED

☐

Changed a file from non-ASCII to ASCII

☐

Changed the margins in cases where the sequence text was "wrapped" down to the next line.

☐

Edited a format error in the Current Application Data section, specifically:

☐

Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_

☐

Added the mandatory heading and subheadings for "Current Application Data".

☐

Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.

☐

Changed the spelling of a mandatory field (the headings or subheadings), specifically:

☐

Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were:

☐

Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

☐

Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐

Inserted colons after headings/subheadings. Headings edited included:

☐

Deleted extra, invalid, headings used by an applicant, specifically:

☒

Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_

☐

Inserted mandatory headings, specifically: \_\_\_\_\_

☐

Corrected an obvious error in the response, specifically: \_\_\_\_\_

☐

Edited identifiers where upper case is used but lower case is required, or vice versa.

☐

Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_

☐

A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.

☐

Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_

☒

Other:

corrected 41407, 41417. numeric identifier

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/580,803

DATE: 07/17/2001

TIME: 14:59:28

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07172001\I580803.raw

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3 <110> APPLICANT: KLAGSBRUN, Michael
4     SOKER, Shay
5     MIAO, Hua Quan
7 <120> TITLE OF INVENTION: ANTAGONISTS OF NEUROPILIN RECEPTOR FUNCTION AND USE THEREOF
9 <130> FILE REFERENCE: 48802 C
11 <140> CURRENT APPLICATION NUMBER: 09/580,803
12 <141> CURRENT FILING DATE: 2000-05-30
14 <150> PRIOR APPLICATION NUMBER: 60/069,155
15 <151> PRIOR FILING DATE: 1997-12-09
17 <150> PRIOR APPLICATION NUMBER: 60/069,687
18 <151> PRIOR FILING DATE: 1997-12-29
20 <150> PRIOR APPLICATION NUMBER: 60/078,541
21 <151> PRIOR FILING DATE: 1998-03-19
24 <160> NUMBER OF SEQ ID NOS: 11
26 <170> SOFTWARE: FastSEQ for Windows Version 3.0
28 <210> SEQ ID NO: 1
29 <211> LENGTH: 5653
30 <212> TYPE: DNA
31 <213> ORGANISM: human
33 <400> SEQUENCE: 1
34 aagggagagg aagccggagc taaatgacag gatgcaggcg acttgagaca caaaaagaga      60
35 agcgttcctc tcggatccag gcattgcctc gctgctttct tttctccaag acgggctgag      120
36 gattgtacag ctctaggcgg agttggggct ctctggatcg cttagattct cctctttgct      180
37 gcatttcccc ccagtcctc gttctccgcg gtctgcctgc ggaccggag aagggagaat      240
38 ggagaggggg tctgcgctcc tctgcgccgt gctgcctc gtcctcgccc cggccggcgc      300
39 ttttcgcaac gataaatgtg gcgatactat aaaaattgaa agccccgggt accttacatc      360
40 tcctggttat cctcattctt atcacccaag tgaaaaatgc gaatggctga ttcaggctcc      420
41 ggaccatac cagagaatta tgatcaactt caaccctcac ttcgatttgg aggacagaga      480
42 ctgcaagtat gactacgtgg aagtgttcga tggagaaaat gaaaatggac attttagggg      540
43 aaagttctgt ggaaagatag cccctcctcc tgttgtgtct tcagggccat ttctttttat      600
44 caaatttgtc tctgactacg aaacacatgg tgcaggattt tccatacgtt atgaaatttt      660
45 caagagaggt cctgaatgtt cccagaacta cacaacacct agtggagtga taaagtcccc      720
46 cggattccct gaaaaatatc ccaacagcct tgaatgcact tatattgtct ttgcgcaaaa      780
47 gatgtcagag attatcctgg aatttgaaa ctttgacctg gagcctgact caaatcctcc      840
48 aggggggatg ttctgtcgct acgaccggct agaaatctgg gatggattcc ctgatgttgg      900
49 ccctcacatt gggcgttact gtggacagaa aacaccaggt cgaatccgat cctcatcggg      960
50 cattctctcc atggtttttt acaccgacag cgcgatagca aaagaaggtt tctcagcaaa     1020
51 ctacagtgtc ttgcagagca gtgtctcaga agatttcaaa tgtatggaag ctctgggcat     1080
52 ggaatcagga gaaattcatt ctgaccagat cacagcttct tcccagtata gcaccaactg     1140
53 gtctgcagag cgctcccgcc tgaactaccc tgagaatggg tggactcccc gagaggattc     1200
54 ctaccgagag tggatacagg tagacttggg cttctctgcg tttgtcacgg ctgtcgggac     1260
55 acagggcgcc atttcaaaa aaaccaagaa gaaatattat gtcaagactt acaagatcga     1320
56 cgtagctcc aacggggaag actggtcac cataaaagaa ggaaacaaac ctgttctctt     1380
57 tcagggaac accaaccaca cagatgttgt ggttcagta ttcccaaac cactgataac     1440
58 tcgatttgtc cgaatcaagc ctgcaacttg ggaaactggc atatctatga gatttgaagt     1500
59 atacggttgc aagataacag attatccttg ctctggaatg ttgggtatgg tgtctggact     1560
60 tatttctgac tcccagatca catcatccaa ccaaggggac agaaactgga tgctgaaaa     1620

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61	catccgcctg	gtaaccagtc	gctctggctg	ggcacttcca	cccgacctc	attcctacat	1680
62	caatgagtg	ctccaaatag	acctggggga	ggagaagatc	gtgaggggca	tcatcattca	1740
63	gggtgggaag	caccgagaga	acaaggtgtt	catgaggaag	ttcaagatcg	ggtacagcaa	1800
64	caacggctcg	gactggaaga	tgatcatgga	tgacagcaaa	cgcaaggoga	agtcttttga	1860
65	gggcaacaac	aactatgata	cacctgagct	gcggaactttt	ccagctctct	ccacgcgatt	1920
66	catcaggatc	taccccgaga	gagccactca	tggcggaactg	gggctcagaa	tggagctgct	1980
67	gggctgtgaa	gtggaagccc	ctacagctgg	accgaccact	cccaacggga	acttggtgga	2040
68	tgaatgtgat	gacgaccagg	ccaactgcca	cagtggaaaca	ggtgatgact	tccagctcac	2100
69	aggtggcacc	actgtgctgg	ccacagaaaa	gcccacggtc	atagacagca	ccatacaatc	2160
70	agagtttcca	acatatggtt	ttaactgtga	atltggctgg	ggctctcaca	agaccttctg	2220
71	ccactgggaa	catgacaatc	acgtgcagct	caagtggagt	gtgttgacca	gcaagacggg	2280
72	acccattcag	gatcacacag	gagatggcaa	cttcatctat	tcccaagctg	acgaaaatca	2340
73	gaagggcaaa	gtggctcgcc	tggtagccc	tgtggtttat	tcccagaact	ctgcccactg	2400
74	catgaccttc	tggatatcaca	tgtctgggtc	ccacgtoggc	acactcaggg	tcaaactgcg	2460
75	ctaccagaag	ccagaggagt	acgatcagct	ggtctggatg	gccattggac	accaaggtga	2520
76	ccactggaag	gaagggcgtg	tcttgcctca	caagtctctg	aaactttatc	aggtgatttt	2580
77	cgagggcgaa	atcggaagaag	gaaaccttgg	tgggattgct	gtggatgaca	ttagtattaa	2640
78	caaccacatt	tcacaagaag	attgtgcaaa	accagcagac	ctggataaaa	agaaccacaga	2700
79	aattaaaatt	gatgaaacag	ggagcacgcc	aggatacgaa	ggtgaaggag	aaggtgacaa	2760
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82	tgcctgttgg	cataatggga	tgtcagaaag	aaacttgtct	gccctggaga	actataactt	2940
83	tgaacttgtg	gatggtgtga	agttgaaaaa	agacaaactg	aatacacaga	gtacttattc	3000
84	ggaggcatga	aggcagacag	agatgaaaag	acagtcaaag	gacggaagtg	gaaggacggg	3060
85	agtgaactgg	ggagctgttg	atctttcaat	atacaggctg	ggaagtgtgt	tgatgaccac	3120
86	tgagccaggc	ttttctcagg	agcttcaatg	agtatggccg	acagacatgg	acaaggagct	3180
87	gtgttcacca	tggactcat	gtgcagtcag	cttttttctt	gttggtttca	tttgaataat	3240
88	cagatgctgg	tgttgagacc	aagtatgatt	gacataatca	ttcatttcga	cccctcctgc	3300
89	ccctctctct	ctctctctct	tcccctttgt	ggattctttt	tggaaactga	gcgaaatcca	3360
90	agatgctggc	accaagcgta	ttccgtgtgg	ccctttggat	ggacatgcta	cctgaaaccc	3420
91	agtgcccgag	atatactaga	atcaccgcag	ttcagtgga	tcctgaagtt	gtacttgtgt	3480
92	ataattgccc	gcgtcgtgca	taggcaaaaga	aggattaggg	tgttttcttt	ttaaagtact	3540
93	gtagcctcag	tactggtgta	gtgtgtcagc	tctgtttacg	aagcaatact	gtccagtttt	3600
94	cttgcctgtt	ttccggtgtt	gtactaaacc	tcgtgcttgt	gaactccata	cagaaaacgg	3660
95	tgccatccct	gaacacggct	ggccactggg	tatactgctg	acaaccgcaa	caacaaaaac	3720
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97	cattgtgtta	cattaacgac	ccactctgct	tcttgcctgg	gaaagccctg	ctctttaatc	3840
98	aaactctggt	ggcccaactg	ctaagaagaa	agtttatatt	cgtgtgagat	gccagccctt	3900
99	ccgggcaggc	aagggtctct	aagatttggc	aacgtggctt	aattgttctg	ctttttctgt	3960
100	agttcaattt	catgtttctt	gaccttttgg	tataaagcta	caatattctc	tcttattgtt	4020
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102	tgtctttttt	ctctcttaga	attggaggat	ttgccattgt	ccaggaaaaga	aacttgcagc	4140
103	tttaacctgc	tgggaatggc	aaacgatttt	actagacttt	atgtttaaaa	ataaataaat	4200
104	aagggaattt	cctaactttg	ccctccaaag	tctaactttg	gttttcttgt	taactggtta	4260
105	aagtgcagct	atcttttttc	cttatctatt	ctattcaaaa	tgacctttga	tagaaatgtt	4320
106	ggcatttagt	agaaatagtg	ataagttgag	gaaagaaata	atacaaattg	gctttcaagt	4380
107	gagacccaaa	ggaagaactg	gataaaatct	tccaaatcca	aaagcatgag	atltttctat	4440
108	ccaaatatgc	aaaaatgacc	caagagaact	ttcttatatt	gctactgagt	cacacaaggg	4500
109	aagtggaagg	aagaacagtt	aatttaagaa	tgaactata	aatcctgatg	cctgggggtc	4560

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/580,803

DATE: 07/17/2001

TIME: 14:59:28

Input Set : A:\Pto.amc

Output Set: N:\CRF3\07172001\I580803.raw

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110 aagtatttta agataagagg gggaaaaaca cataaagtca aacaaatggt ttaaaaattc 4620
111 ataacagcaa ccttgaaaaa atagacttaa atgaatgctt ctagaaactt ccagcggctc 4680
112 acaaagaata agcctgcctt agggctggca acatctaagc ctctaacagc acagggaagc 4740
113 aaatatctta ccaggcagcc tatgaattaa cccaaagaag ctttggttg ttttggtgga 4800
114 tttttatcat gccatggttg acatgagatt ttttagatct tccttcccca cattgctaga 4860
115 cgtctcactc aaagacattt gttgggagtc acatttgcac catagacgag acagtccatt 4920
116 catcttagtt aaattggatt gagaatgcct tttgtttcca ggaaaatatt gatcaccatg 4980
117 aaagaagaat agttttttgt ccccagagac attcatttag ttgatataat cctaccagaa 5040
118 ggaaagcact aagaaacact cgtttgttgt ttttaaaggc aacagactta aagttgtcct 5100
119 cagccaagga aaaatgatac tgcaacttta aaatttaaag tatcttgacac tgataaatat 5160
120 atttaaaaaat tatatgttta taaagttatt aatttgtaaa ggcagtgtta caaaatgttc 5220
121 agtttatatt gtttttagatt gttttgtaat ttttaaaggc gtaaaataac atataaatat 5280
122 atttaaaaaat tatatgttta taaagttatt aatttgtaaa ggcagtgtta caaaatgttc 5340
123 agtttatatt gtttttagatt gttttgtaat ttttaaaggc gtaaaataac atattttttc 5400
124 tttatgaaa tctataaaac tttctgtagt aaaatgtttt cattttactg gtatattatt 5460
125 gcttcatggt ttgtaccatc ataagatttt gtgcagattt tttttacaga aattattatt 5520
126 ttctatgaca atatgacact tgtaaattgt tgtttcaaaa tgaacagcga agccttaact 5580
127 ttaaatgaca tttgtattct cagacactga gtagcataaa aaccacatga actgaactgt 5640
128 aacttaaatt ctt 5653

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132 &lt;210&gt; SEQ ID NO: 2

133 &lt;211&gt; LENGTH: 923

134 &lt;212&gt; TYPE: PRT

135 &lt;213&gt; ORGANISM: human

137 &lt;400&gt; SEQUENCE: 2

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138 Met Glu Arg Gly Leu Pro Leu Leu Cys Ala Val Leu Ala Leu Val Leu
139 1 5 10 15
140 Ala Pro Ala Gly Ala Phe Arg Asn Asp Lys Cys Gly Asp Thr Ile Lys
141 20 25 30
142 Ile Glu Ser Pro Gly Tyr Leu Thr Ser Pro Gly Tyr Pro His Ser Tyr
143 35 40 45
144 His Pro Ser Glu Lys Cys Glu Trp Leu Ile Gln Ala Pro Asp Pro Tyr
145 50 55 60
146 Gln Arg Ile Met Ile Asn Phe Asn Pro His Phe Asp Leu Glu Asp Arg
147 65 70 75 80
148 Asp Cys Lys Tyr Asp Tyr Val Glu Val Phe Asp Gly Glu Asn Glu Asn
149 85 90 95
150 Gly His Phe Arg Gly Lys Phe Cys Gly Lys Ile Ala Pro Pro Val
151 100 105 110
152 Val Ser Ser Gly Pro Phe Leu Phe Ile Lys Phe Val Ser Asp Tyr Glu
153 115 120 125
154 Thr His Gly Ala Gly Phe Ser Ile Arg Tyr Glu Ile Phe Lys Arg Gly
155 130 135 140
156 Pro Glu Cys Ser Gln Asn Tyr Thr Thr Pro Ser Gly Val Ile Lys Ser
157 145 150 155 160
158 Pro Gly Phe Pro Glu Lys Tyr Pro Asn Ser Leu Glu Cys Thr Tyr Ile
159 165 170 175
160 Val Phe Ala Pro Lys Met Ser Glu Ile Ile Leu Glu Phe Glu Ser Phe
161 180 185 190
162 Asp Leu Glu Pro Asp Ser Asn Pro Pro Gly Gly Met Phe Cys Arg Tyr

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163		195		200		205											
164	Asp	Arg	Leu	Glu	Ile	Trp	Asp	Gly	Phe	Pro	Asp	Val	Gly	Pro	His	Ile	
165		210					215				220						
166	Gly	Arg	Tyr	Cys	Gly	Gln	Lys	Thr	Pro	Gly	Arg	Ile	Arg	Ser	Ser	Ser	
167	225					230					235					240	
168	Gly	Ile	Leu	Ser	Met	Val	Phe	Tyr	Thr	Asp	Ser	Ala	Ile	Ala	Lys	Glu	
169				245						250					255		
170	Gly	Phe	Ser	Ala	Asn	Tyr	Ser	Val	Leu	Gln	Ser	Ser	Val	Ser	Glu	Asp	
171				260					265					270			
172	Phe	Lys	Cys	Met	Glu	Ala	Leu	Gly	Met	Glu	Ser	Gly	Glu	Ile	His	Ser	
173			275					280					285				
174	Asp	Gln	Ile	Thr	Ala	Ser	Ser	Gln	Tyr	Ser	Thr	Asn	Trp	Ser	Ala	Glu	
175		290					295				300						
176	Arg	Ser	Arg	Leu	Asn	Tyr	Pro	Glu	Asn	Gly	Trp	Thr	Pro	Gly	Glu	Asp	
177	305					310				315						320	
178	Ser	Tyr	Arg	Glu	Trp	Ile	Gln	Val	Asp	Leu	Gly	Leu	Leu	Arg	Phe	Val	
179				325						330					335		
180	Thr	Ala	Val	Gly	Thr	Gln	Gly	Ala	Ile	Ser	Lys	Glu	Thr	Lys	Lys	Lys	
181			340					345						350			
182	Tyr	Tyr	Val	Lys	Thr	Tyr	Lys	Ile	Asp	Val	Ser	Ser	Asn	Gly	Glu	Asp	
183			355				360						365				
184	Trp	Ile	Thr	Ile	Lys	Glu	Gly	Asn	Lys	Pro	Val	Leu	Phe	Gln	Gly	Asn	
185		370					375				380						
186	Thr	Asn	Pro	Thr	Asp	Val	Val	Val	Ala	Val	Phe	Pro	Lys	Pro	Leu	Ile	
187	385					390				395						400	
188	Thr	Arg	Phe	Val	Arg	Ile	Lys	Pro	Ala	Thr	Trp	Glu	Thr	Gly	Ile	Ser	
189				405					410						415		
190	Met	Arg	Phe	Glu	Val	Tyr	Gly	Cys	Lys	Ile	Thr	Asp	Tyr	Pro	Cys	Ser	
191			420				425						430				
192	Gly	Met	Leu	Gly	Met	Val	Ser	Gly	Leu	Ile	Ser	Asp	Ser	Gln	Ile	Thr	
193			435				440					445					
194	Ser	Ser	Asn	Gln	Gly	Asp	Arg	Asn	Trp	Met	Pro	Glu	Asn	Ile	Arg	Leu	
195		450					455				460						
196	Val	Thr	Ser	Arg	Ser	Gly	Trp	Ala	Leu	Pro	Pro	Ala	Pro	His	Ser	Tyr	
197	465					470				475						480	
198	Ile	Asn	Glu	Trp	Leu	Gln	Ile	Asp	Leu	Gly	Glu	Glu	Lys	Ile	Val	Arg	
199				485					490						495		
200	Gly	Ile	Ile	Ile	Gln	Gly	Gly	Lys	His	Arg	Glu	Asn	Lys	Val	Phe	Met	
201			500					505						510			
202	Arg	Lys	Phe	Lys	Ile	Gly	Tyr	Ser	Asn	Asn	Gly	Ser	Asp	Trp	Lys	Met	
203			515					520					525				
204	Ile	Met	Asp	Asp	Ser	Lys	Arg	Lys	Ala	Lys	Ser	Phe	Glu	Gly	Asn	Asn	
205		530					535				540						
206	Asn	Tyr	Asp	Thr	Pro	Glu	Leu	Arg	Thr	Phe	Pro	Ala	Leu	Ser	Thr	Arg	
207	545					550				555						560	
208	Phe	Ile	Arg	Ile	Tyr	Pro	Glu	Arg	Ala	Thr	His	Gly	Gly	Leu	Gly	Leu	
209				565					570						575		
210	Arg	Met	Glu	Leu	Leu	Gly	Cys	Glu	Val	Glu	Ala	Pro	Thr	Ala	Gly	Pro	
211			580					585						590			

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212 Thr Thr Pro Asn Gly Asn Leu Val Asp Glu Cys Asp Asp Asp Gln Ala
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214 Asn Cys His Ser Gly Thr Gly Asp Asp Phe Gln Leu Thr Gly Gly Thr
215          610                      615                      620
216 Thr Val Leu Ala Thr Glu Lys Pro Thr Val Ile Asp Ser Thr Ile Gln
217          625                      630                      635                      640
218 Ser Glu Phe Pro Thr Tyr Gly Phe Asn Cys Glu Phe Gly Trp Gly Ser
219          645                      650                      655
220 His Lys Thr Phe Cys His Trp Glu His Asp Asn His Val Gln Leu Lys
221          660                      665                      670
222 Trp Ser Val Leu Thr Ser Lys Thr Gly Pro Ile Gln Asp His Thr Gly
223          675                      680                      685
224 Asp Gly Asn Phe Ile Tyr Ser Gln Ala Asp Glu Asn Gln Lys Gly Lys
225          690                      695                      700
226 Val Ala Arg Leu Val Ser Pro Val Val Tyr Ser Gln Asn Ser Ala His
227          705                      710                      715                      720
228 Cys Met Thr Phe Trp Tyr His Met Ser Gly Ser His Val Gly Thr Leu
229          725                      730                      735
230 Arg Val Lys Leu Arg Tyr Gln Lys Pro Glu Glu Tyr Asp Gln Leu Val
231          740                      745                      750
232 Trp Met Ala Ile Gly His Gln Gly Asp His Trp Lys Glu Gly Arg Val
233          755                      760                      765
234 Leu Leu His Lys Ser Leu Lys Leu Tyr Gln Val Ile Phe Glu Gly Glu
235          770                      775                      780
236 Ile Gly Lys Gly Asn Leu Gly Gly Ile Ala Val Asp Asp Ile Ser Ile
237          785                      790                      795                      800
238 Asn Asn His Ile Ser Gln Glu Asp Cys Ala Lys Pro Ala Asp Leu Asp
239          805                      810                      815
240 Lys Lys Asn Pro Glu Ile Lys Ile Asp Glu Thr Gly Ser Thr Pro Gly
241          820                      825                      830
242 Tyr Glu Gly Glu Gly Glu Gly Asp Lys Asn Ile Ser Arg Lys Pro Gly
243          835                      840                      845
244 Asn Val Leu Lys Thr Leu Asp Pro Ile Leu Ile Thr Ile Ile Ala Met
245          850                      855                      860
246 Ser Ala Leu Gly Val Leu Leu Gly Ala Val Cys Gly Val Val Leu Tyr
247          865                      870                      875                      880
248 Cys Ala Cys Trp His Asn Gly Met Ser Glu Arg Asn Leu Ser Ala Leu
249          885                      890                      895
250 Glu Asn Tyr Asn Phe Glu Leu Val Asp Gly Val Lys Leu Lys Lys Asp
251          900                      905                      910
252 Lys Leu Asn Thr Gln Ser Thr Tyr Ser Glu Ala
253          915                      920
256 <210> SEQ ID NO: 3
257 <211> LENGTH: 3404
258 <212> TYPE: DNA
259 <213> ORGANISM: human
261 <400> SEQUENCE: 3
262 gaattcggca cgaggggaaa ataaaagaga gaaaaacaca aagatttaaa caagaaacct      60
263 acgaaccag ctctggaaag agccaccttc tccaaaatgg atatgtttcc tctcacctgg      120

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**VERIFICATION SUMMARY**

**PATENT APPLICATION: US/09/580,803**

**DATE: 07/17/2001**

**TIME: 14:59:29**

**Input Set : A:\Pto.amc**

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1642

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/580,803

DATE: 07/17/2001

TIME: 14:23:03

Input Set : A:\ES.txt

Output Set: N:\CRF3\07172001\I580803.raw

Does Not Comply  
Corrected Diskette Needed

3 <110> APPLICANT: KLAGSBRUN, Michael  
 4 SOKER, Shay  
 5 MIAO, Hua Quan  
 7 <120> TITLE OF INVENTION: ANTAGONISTS OF NEUROPILIN RECEPTOR FUNCTION AND USE THEREOF  
 9 <130> FILE REFERENCE: 48802 C  
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/580,803  
 C--> 11 <141> CURRENT FILING DATE: 2000-05-30  
 11 ~~<150> PRIOR APPLICATION NUMBER: 09/580,803~~  
 12 ~~<151> PRIOR FILING DATE: 2000-05-30~~  
 14 <150> PRIOR APPLICATION NUMBER: 60/069,155  
 15 <151> PRIOR FILING DATE: 1997-12-09  
 17 <150> PRIOR APPLICATION NUMBER: 60/069,687  
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 20 <150> PRIOR APPLICATION NUMBER: 60/078,541  
 21 <151> PRIOR FILING DATE: 1998-03-19  
 24 <160> NUMBER OF SEQ ID NOS: 11  
 26 <170> SOFTWARE: FastSEQ for Windows Version 3.0

## ERRORED SEQUENCES

500 <210> SEQ ID NO: 11  
 501 <211> LENGTH: 44  
 502 <212> TYPE: PRT  
 503 <213> ORGANISM: human  
 505 <400> SEQUENCE: 11  
 506 Pro Cys Gly Pro Cys Ser Glu Arg Arg Lys His Leu Phe Val Gln Asp  
 507 1 5 10 15  
 508 Pro Gln Thr Cys Lys Cys Ser Cys Lys Asn Thr Asp Ser Arg Cys Lys  
 509 20 25 30  
 510 Ala Arg Gln Leu Glu Leu Asn Glu Arg Thr Cys Arg  
 511 35 40  
 E--> 515 (Footnote continued from previous page)  
 E--> 516 (Footnote continued on next page)



## VERIFICATION SUMMARY

PATENT APPLICATION: US/09/580,803

DATE: 07/17/2001

TIME: 14:23:04

Input Set : A:\ES.txt

Output Set: N:\CRF3\07172001\I580803.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No  
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:515 M:333 E: Wrong sequence grouping, Amino acids not in groups!  
L:515 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:5  
L:516 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:11  
L:516 M:333 E: Wrong sequence grouping, Amino acids not in groups!  
L:516 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:5  
L:516 M:252 E: No. of Seq. differs, <211>LENGTH:Input:44 Found:54 SEQ:11